



OIPE

RAW SEQUENCE LISTING

DATE: 02/27/2002

PATENT APPLICATION: US/09/911,047

TIME: 10:55:08

Input Set : A:\E104720060-SeqLst-072301.txt

Output Set: N:\CRF3\02272002\I911047.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Erikson, Glen
 4 Daksis, Jasmine
 5 Picard, Pierre
 7 <120> TITLE OF INVENTION: HOMOGENEOUS ASSAY OF BIOPOLYMER BINDING BY
 8 MEANS OF MULTIPLE MEASUREMENTS UNDER VARIED CONDITIONS
 10 <130> FILE REFERENCE: E1047/20060
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/911,047
 C--> 13 <141> CURRENT FILING DATE: 2001-07-23
 15 <160> NUMBER OF SEQ ID NOS: 9
 17 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

125 <210> SEQ ID NO: 9
 126 <211> LENGTH: 15
 127 <212> TYPE: DNA
 128 <213> ORGANISM: Artificial Sequence
 130 <220> FEATURE:
 131 <223> OTHER INFORMATION: Description of Artificial Sequence: ssDNA probe wherein the
 3' end of each base is covalently bonded to a lysine N-terminal leaving a free carboxyl group
 133 <400> SEQUENCE: 9
 E--> 134 tatagtagaa accac 15←

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/911,047

DATE: 02/27/2002

TIME: 10:55:09

Input Set : A:\E104720060-SeqLst-072301.txt

Output Set: N:\CRF3\02272002\I911047.raw

L:12 M:270 C: Current Application Number differs, Replaced Application Number
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:121 M:283 W: Missing Blank Line separator, <400> field identifier
L:134 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:15 SEQ:9



OIIPE

RAW SEQUENCE LISTING

DATE: 02/27/2002

PATENT APPLICATION: US/09/911,047

TIME: 10:55:46

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\02272002\I911047.raw

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3 <110> APPLICANT: Erikson, Glen
4   Daksis, Jasmine
5   Picard, Pierre
7 <120> TITLE OF INVENTION: HOMOGENEOUS ASSAY OF BIOPOLYMER BINDING BY
8   MEANS OF MULTIPLE MEASUREMENTS UNDER VARIED CONDITIONS
10 <130> FILE REFERENCE: E1047/20060
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/911,047
C--> 13 <141> CURRENT FILING DATE: 2001-07-23
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RAW SEQUENCE LISTING

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Input Set : A:\PTO.AMC.txt

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3' end of each
133      base is covalently bonded to a lysine N-terminal leaving a free carboxyl group
135 <400> SEQUENCE: 9
136 tatagtagaa accac 15

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/911,047

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

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